-FORM PTO-1449 (REV. 7-85)

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ATTY. DOCKET NO. 60063USDIV2 APPLICATION NO. 10/623,880 APPLICANT BECK, J. and BARNETT, J.

FILING DATE July 21, 2003

Group

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U.S. PATENT DOCUMENTS

PADEMARK		U.S. PATENT DOCUMENTS					
EXAMINER INITIAL		DOCUMENT NUMBER	DATE	NAME	CLASS	SUBCLASS	FILING DATE
1	AA	4,683,202	7/28/87	Kary B. Mullis	-435	91	10/25/85
	AB	4,683,195	7/28/87	Mullis, et al.	435	6_	2/7/86
	AC	5,585,238	12/17/96	James. Ligon and James. Beck	435	6	4/25/94
	AD	5,800,997	9/1/98	James J. Beck	435	6	11/1/96
1	AE	5,955,274	9/21/99	James Ligon and James Beck	435	6	4/19/95

OTHER DOCUMENTS (Including Author, Title, Date, Pertinent pages, Etc.)

M	AA	Adaskaveg, J.E. and Hartin, R.J., Characterization of Colletotrichum acutatum Isolates Causing Anthracnose of Almond and Peach in California Etiology, Vol. 87, No. 9 (1997), pp. 979-987
	АВ	Barker, I., et al. "Strawberry Blackspot Disease (Colletotrichum acutatum)" in Schots A., Dewey FM, Oliver R. (Eds.). Modern assay for plant pathogenic fungi: identification, detection and quantification (Wallingford, Oxford: CAB International, 1994), pp. 179-182.
·	AC	Förster, H. and Adaskaveg, J.E., Identification of Subpopulations of Colletotrichum acutatum and Epidemiology of Almond Anthracnose in California Phytopathology, Vol. 89, No. 11 (1999), pp. 1056-1065
	AD	Johanson, A. and Jeger M. Use of PCR for detection of Mycosphaerella fijiensis and M. musicola, the causal agents of Sigatoka leaf spots in banana and plantain Mycological Research, Vol. 97, No. 6 (1993), pp. 670-674.
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	AF	Lee, S.B. and Taylor, J.W., "Isolation of DNA from fungal mycelia and single spores." In: eds. Innis, et al., <i>PCR Protocols: A Guide to Methods and Applications</i> (New York, Academic Press, Inc., 1990) Pp. 282-287.
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	АН	Nazar, et al. Potential use of PCR-amplified ribosomal intergenic sequences in the detection and differentiation of verticillium wilt pathogens Physiological and Molecular Plant Pathology, Vol. 39, (1991), pp. 1-11.
V	Al	Pryor, B.M. and Gilbertson, R.L. Molecular phylogenetic relationships amongst Alternaria species and related fungi based upon analysis of nuclear ITS and mt SSU rDNA sequences Mycological Research, Vol. 104, Part 11 (Nov. 2000), pp. 1312-1321
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M	AK	Schesser, K., et al. Use of Polymerase Chain Reaction to Detect the Take-All Fungus, Gaeumannomyces graminis, in Infected Wheat Plants Applied and Environmental Microbiology, Vol. 57, No. 2 (1990), pp. 553-556
	AL	Schnabel, G. et al. Characterization of Ribosomal DNA from Venturia inaequalis and Its Phylogenetic Relationship to rDNA from Other Tree-Fruit Venturia Species Phytopathology, Vol. 89, No. 1 (1999), pp. 100-108
	AM	Sreenivasaprasad, S., et al. Phylogeny and systematics of 18 Colletotrichum species based on ribosomal DNA spacer sequences Genome, Vol. 39 (1996), pp. 499-512
	AN	Teviotdale, B.L., et al. First Report of Alternaria Leaf Spot of Almond Caused by Species in the Alternaria alternata Complex in California Plant Disease, Vol. 85, No. 5 (May, 2001), pp. 558
		Wang et al, PCR amplification from single seeds, facilitating DNA marker-assisted breeding Nucleic Acids Research, Vol. 21, No. 10 (1993), pp. 2527
	AO	White, et al. "Amplification and direct sequencing of fungal ribosomal RNA genes for phylogenetics." In: eds. Innis, et al., <i>PCR Protocols: A Guide to Methods and Applications</i> (New York, Academic Press, Inc., 1990) Pp. 315-322.
	AP	Zur, et al. Development of a Polymerase Chain Reaction-Based Assay for the Detection of Alternaria Fungal Contamination in Food Products Journal of Food Protection, Vol. 62, No. 10 (1999), pp. 1191-1197
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M	АХ	GenBank Accession Number AF229460 [online], retrieved on 2001-10-26]. Retrieved from the Internet: <url: http:="" www.ncbi.nlm.nih.gov=""></url:>
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